Serial Number: _09/591,737_   **ENTERED**

- ☐ Changed a file from non-ASCII to ASCII

- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

- ☐ Edited a format error in the Current Application Data section, specifically:
  _____

- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

- ☐ Added the mandatory heading and subheadings for "Current Application Data".

- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
  _____

- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
  _____

- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
  _____

- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

- ☐ Inserted colons after headings/subheadings. Headings edited included:
  _____

- ☐ Deleted extra, invalid, headings used by an applicant, specifically:
  _____

- ☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____.

- ☐ Inserted mandatory headings, specifically: _____

- ☐ Corrected an obvious error in the response, specifically:
  _____

- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

- ☐ Corrected an error in the Number of Sequences field, specifically:
  _____

- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

- ☐ Other:
  _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.   3/1/95

> **This Raw Listing contains the General Information Section and up to first 5 pages.**

```
 1   <110> APPLICANT: Curiel et al.
 2   <120> TITLE OF INVENTION: Immunomodulation by genetic modification of
 3         dendritic cells and B cells
 4   <130> FILE REFERENCE: D6167CIP
 5   <140> CURRENT APPLICATION NUMBER: US/09/591,737
 6   <141> CURRENT FILING DATE: 2000-06-12
 7   <150> EARLIER APPLICATION NUMBER: 60/102,257
 8   <151> EARLIER FILING DATE: 1998-09-29
 9   <160> NUMBER OF SEQ ID NOS: 1
10   <210> SEQ ID NO 1
11   <211> LENGTH: 9
12   <212> TYPE: PRT
13   <213> ORGANISM: Unknown
14   <220> FEATURE:
15   <221> NAME/KEY: DOMAIN
16   <222> LOCATION:
17   <223> OTHER INFORMATION: RGD motif that binds specifically to _v_3 and
18         _v_5 integrins
19   <400> SEQUENCE: 1
20         Cys Asp Cys Arg Gly Asp Cys Phe Cys
21                         5
```

Line ? Error/Warning   Original Text
----- - --- ---

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION **US/09/591,737**

DATE: 06/21/2000
TIME: 11:05:33

Input Set: **I591737.RAW**

PREVIOUSLY ERRORED SEQUENCES-EDITED

```
  1     <210> 1
  2     <211> 9
  3     <212> PRT
  4     <213> Unknown
  5     <220>
  6     <221> DOMAIN
  7     <222>
  8     <223> RGD motif that binds specifically to _v_3 and
  9           _v_5 integrins
 10     <400> 1
 11           Cys Asp Cys Arg Gly Asp Cys Phe Cys
 12                           5
```

+--+
| **This Raw Listing contains the General** |
| **Information Section and those Sequences** |
| **containing ERRORS.** |
+--+

    1   <110> Curiel et al.
    2   <120> Immunomodulation by genetic modification of
    3   dendritic cells and B cells
    4   <130> D6167CIP
    5   <140> US/09/591,737
    6   <141> 2000-06-12
    7   <150> 60/102,257
    8   <151> 1998-09-29
    9   <160> 1

ERRORED SEQUENCES FOLLOW

---

          10   <210> 1
E-->   11   **<211> 9**
          12   <212> PRT
          13   <213> Unknown
          14   <220>
          15   <221> DOMAIN
          16   <222>
          17   <223> RGD motif that binds specifically to _v_3 and
          18   _v_5 integrins
          19   <400> 1
          20   Cys Asp Cys Arg Gly Asp Cys Phe Cys
          21   5
E-->   22   SEQ

```
Line  ? Error/Warning                       Original Text
----- - ------------------------------------ ------------------------------------------
   11 E Input 9, Calc Seq.Length 10 differ    <211>  9
   22 E Wrong Amino Acid Designator           SEQ 1
```